

Please replace the paragraph beginning on page 6, line 26 - page 7, line 4 with the following:

B | sVpr¹⁻⁹⁶:

H - Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn - Glu - Trp - Thr - Leu - Glu - Leu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe - Pro - Arg - Ile - Trp - Leu - His - Asn - Leu - Gly - Gln - His - Ile - Tyr - Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln - Gln - Leu - Leu - Phe - Ile - His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg - Ile - Gly - Val - Thr - Arg - Gln - Arg - Arg - Ala - Arg - Asn - Gly - Ala - Ser - Arg - Ser-OH (SEQ ID NO: 1)

Please replace the paragraph on page 7, lines 6-10 with the following:

B | sVpr¹⁻⁴⁷:

H-Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn - Glu - Trp - Thr - Leu - Glu - Leu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe - Pro - Arg - Ile - Trp - Leu - His - Asn - Leu - Gly - Gln - His - Ile - Tyr-NH₂ (SEQ ID NO: 2)

Please replace the paragraph on page 7, lines 12-16 with the following:

B | sVpr⁴⁸⁻⁹⁶:

Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln - Gln - Leu - Leu - Phe - Ile - His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg - Ile - Gly - Val - Thr - Arg - Gln - Arg - Arg - Ala - Arg - Asn - Gly - Ala - Ser - Arg - Ser-OH (SEQ ID NO: 3)

Please replace the paragraph on page 7, lines 18-20 with the following:

sVpr¹⁻²⁰ as mutant *sVpr¹⁻²⁰(Asn^{5,10,14})*:

H-Met - Glu - Gln - Ala - Asn - Glu - Asp - Gln - Gly - Asn - Gln - Arg - Glu - Asn - Tyr - Asn -
B4 Glu - Trp - Thr - Leu-NH₂ (SEQ ID NO: 8), and

Please replace the paragraph on page 7, lines 22-24 with the following:

sVpr²¹⁻⁴⁰ as mutant *sVpr²¹⁻⁴⁰(Asn³⁵)*:

H-Glu - Leu - Leu - Glu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe - Asn - Arg - Ile -
Trp - Leu - His-NH₂ (SEQ ID NO: 9),

Please replace the paragraph on page 8, lines 1-3 with the following:

sVpr¹¹⁻²⁵:

H-Gln - Arg - Glu - Pro - Tyr - Asn - Glu - Trp - Thr - Leu - Glu - Leu - Leu - Glu - Glu-NH₂
(SEQ ID NO: 4),

Please replace the paragraph on page 8, lines 5-7 with the following:

sVpr⁴¹⁻⁵⁵:

H-Asn - Leu - Gly - Gln - His - Ile - Tyr - Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala-NH₂ (SEQ
ID NO: 5),

Please replace the paragraph on page 8, lines 9-11 with the following:

B6 sVpr⁴⁶⁻⁶⁰:

H-Ile - Tyr - Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile-NH₂ (SEQ ID NO: 6),

Please replace the paragraph on page 8, lines 13-15 with the following:

B7 sVpr⁵⁶⁻⁷⁰:

H-Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln - Gln - Leu - Leu - Phe - Ile-NH₂ (SEQ ID NO: 7),

Please replace the paragraph on page 8, lines 17-19 with the following:

B8 sVpr⁶⁶⁻⁸⁰:

H-Gln - Leu - Leu - Phe - Ile - His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg-NH₂ (SEQ ID NO: 10),

Please replace the paragraph on page 8, lines 21-23 with the following:

B9 sVpr⁷⁶⁻⁹⁶:

H-Cys - Arg - His - Ser - Arg - Ile - Gly - Val - Thr - Arg - Gln - Arg - Arg - Ala - Arg - Asn - Gly - Ala - Ser - Arg - Ser-OH (SEQ ID NO: 11).

Please replace the paragraph on page 14, lines 9-17 with the following:

molecular weight: calculated: 11378

found: 11381

H - Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn - Glu
- Trp - Thr - Leu - Glu - Leu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe -
Pro - Arg - Ile - Trp - Leu - His - Asn - Leu - Gly - Gln - His - Ile - Tyr - Glu - Thr - Tyr - Gly - Asp
- Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln - Gln - Leu - Leu - Phe - Ile
- His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg - Ile - Gly - Val - Thr - Arg - Gln - Arg -
Arg - Ala - Arg - Asn - Gly - Ala -
Ser - Arg - Ser - OH (SEQ ID NO: 1).

Please replace the paragraph beginning on page 14, line 24 - page 15, line 2 with the following:

Example 4:

*sVpr*¹⁻⁴⁷

in analogy to examples 1 to 3.

molecular weight: calculated: 5728

found: 5728.8

B13 H - Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn -
Glu - Trp - Thr - Leu - Glu - Leu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His -
Phe - Pro - Arg - Ile - Trp - Leu - His - Asn - Leu - Gly - Gln - His - Ile - Tyr - NH₂ (SEQ ID NO:
9).

Please replace the paragraph on page 15, lines 5-11 with the following:

Example 5:

*Vpr*⁴⁸⁻⁹⁶

in analogy to examples 1 to 3.

B14
Contd.

Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln -
Gln - Leu - Leu - Phe - Ile - His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg - Ile - Gly - Val -
Thr - Arg - Gln - Arg - Arg - Ala - Arg - Asn - Gly - Ala - Ser - Arg - Ser - OH. (SEQ ID NO: 3).

Please replace the paragraph on page 15, lines 13-19 with the following:

Example 6:

sVpr¹⁻²⁰

in analogy to examples 1 to 3.

B15
H - Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn -
Glu - Trp - Thr - Leu - NH₂ (SEQ ID NO: 8).

Figure 5: *sVpr¹⁻²⁰* - mass spectrum (% int. and molecular weight) (%Int. 10% = 111

mV[sum=9505 mV].

Please replace the paragraph on page 15, lines 21-25 with the following:

Example 7:

sVpr¹⁻²⁰ (Asn^{5,10,14})

B16
in analogy to examples 1 to 3.

H - Met - Glu - Gln - Ala - Pro - Glu - Asp - Gln - Gly - Pro - Gln - Arg - Glu - Pro - Tyr - Asn -
Glu - Trp - Thr - Leu - NH₂ (SEQ ID NO: 8).

Please replace the paragraph beginning on page 15, line 27 - page 16, line 3 with the
following:

Example 8:

B17
Contd.
sVpr²¹⁻⁴⁰

in analogy to examples 1 to 3.

Wildtype-sequence:

B17
Conrad H - Glu - Leu - Leu - Glu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe - Asn - Arg - Ile
- Trp - Leu - His - NH₂ (SEQ ID NO: 9).

Please replace the paragraph on page 16, lines 6-10 with the following:

Example 9:

sVpr²¹⁻⁴⁰(Asn³⁵)

B18 in analogy to examples 1 to 3.

H - Glu - Leu - Leu - Glu - Glu - Leu - Lys - Ser - Glu - Ala - Val - Arg - His - Phe - Asn - Arg - Ile
- Trp - Leu - His - NH₂ (SEQ ID NO: 9).

Please replace the paragraph on page 16, lines 12-16 with the following:

Example 10:

sVpr¹¹⁻²⁵:

B19 in analogy to examples 1 to 3.

H - Gln - Arg - Glu - Pro - Tyr - Asn - Glu - Trp - Thr - Leu - Glu - Leu - Leu - Glu - Glu - NH₂
(SEQ ID NO: 4).

Please replace the paragraph on page 16, lines 18-22 with the following:

B20 Example 11:

sVpr⁴¹⁻⁵⁵:

in analogy to examples 1 to 3.

H - Asn - Leu - Gly - Gln - His - Ile - Tyr - Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - NH₂
(SEQ ID NO: 5).

Please replace the paragraph on page 16, lines 24-28 with the following:

B21
Example 12:

sVpr⁴⁶⁻⁶⁰:

in analogy to examples 1 to 3.

H - Ile - Tyr - Glu - Thr - Tyr - Gly - Asp - Thr - Trp - Ala - Gly - Val - Glu - Ala - Ile - NH₂ (SEQ ID NO: 6).

Please replace the paragraph on page 17, lines 1-5 with the following:

B22
Example 13:

sVpr⁵⁶⁻⁷⁰:

in analogy to examples 1 to 3.

H - Gly - Val - Glu - Ala - Ile - Ile - Arg - Ile - Leu - Gln - Gln - Leu - Leu - Phe - Ile - NH₂ (SEQ ID NO: 7).

Please replace the paragraph on page 17, lines 7-11 with the following:

B23
Example 14:

sVpr⁶⁶⁻⁸⁰:

in analogy to examples 1 to 3.

H - Gln - Leu - Leu - Phe - Ile - His - Phe - Arg - Ile - Gly - Cys - Arg - His - Ser - Arg - NH₂ (SEQ ID NO: 10).

Please replace the paragraph on page 17, lines 13-17 with the following:

B24
Example 15:

sVpr⁷⁶⁻⁹⁶

in analogy to examples 1 to 3.

H-Cys - Arg - His - Ser - Arg - Ile - Gly - Val - Thr - Arg - Gln - Arg - Arg - Ala - Arg - Asn - Gly - Ala - Ser - Arg - Ser - OH (SEQ ID NO: 11).